	CTCTGGTGTGGAGGGTGTTCATACCCGTCGGAGTAAAACCGGAGTTCTGGTTGTAATGGCACCCTGAGTA
	10 20 30 40 50 60 70
141233	CTCTCGTGTCGAGGGTGTTCATACCCGTCGGAGTAAAACCGGAGTTCTGGTTGTAATGGCACCCTGAGTA 7.0
606977	CICIGGIGIGGAGGGIGITCATACCCGICGGAGIAAAACCGGAGITCIGGIIGIAATGGCACCCIGAGIA 70
000511	
•	
	CGAGIGICIGITCATTGIGGITGICGAAGCCTGAACATTTGGGIATTTGAAGIGAATATCGXGITTCTTC
	80 90 100 110 120 130 140
141233	CCAGTGTCTCATTGTCGTTGTCGAACCCTGAACATTTGCGTATTTGAAGTGAATATCGNGTTTCTTC 140
606977	CGAGTGTCTGTTCATTGTCGTCGTCAACCCTCAACATTTCGCGTATTTCGAACTGAATATCCACTTTTCTTC 140
000577	
	CTTATTTCGAGCACTAAGACTCGCCTGTCGGCTAGCTGAACCGCTTAACCGAGCGTGAGTTGCCTTGCGC
	150 160 170 180 190 200 210
141233	CTTATTTCGAGCACTAAGACTCGCCTGTCGCCTGAGCTGAACCGCTTAACCGAGCGTGAGTTGCCTTTGCCC 2.1.0
606977	CITATITICASCACIASCACIOCCITICOCCIAGCIGAACCCCITIAACCCAGCGIGAGTIGCCITICCC 210
000517	
	GGAAGGTGACGAGTGAGGTATCTGTATTCCGGAGGCGTAAGAGTTCCTCGGCTCGGTCGG
	220 230 240 250 260 270 280
141233	GGAAGGIGAGGAGTGAGGTATCIGIATTCCGGAGGCGTAAGAGTTCCTCGGCTCGG
606977	GGAAGGIGACGAGIGAGGIATCIGIATTCCGGAGGCGIAAGAGITCCTCGGCTCGG
<b>x</b>	
141233	280
606977	280